From the Ground up: Assessing Bacterial Diversity in the Air Space of an Urban Park

Janet L. Pelley

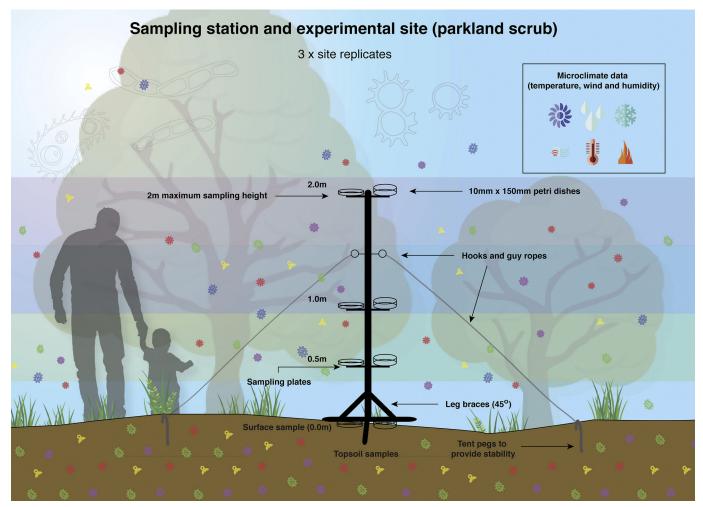
https://doi.org/10.1289/EHP8736

Emerging research has linked exposure to diverse airborne microbial communities to healthy human immune systems. But few studies have examined how communities of airborne microbes (aerobiomes) vary over time and space²; none are known to have investigated whether these communities change with height from the ground. A recent proof-of-concept study in *Environmental Health Perspectives* described how the diversity of airborne bacterial communities decreased from ground level to 2 m in height in an urban park, noting that the makeup of the communities also varied, depending on height.³

"Exposure to a diverse set of microbes from our environment trains our immune systems to respond appropriately to pathogens," says Graham Rook, a professor of medical microbiology at University College London who was not involved in the study. Lack of exposure to bacterial diversity has been associated with autoimmune diseases, allergies, Alzheimer's disease, and inflammatory bowel disease.

Airborne microbial communities are important for human health because our skin and mucous membranes are exposed to them every day. Yet the community structure of the aerobiome remains little known, says lead study author Jake Robinson, a doctoral student in microbial ecology at England's University of Sheffield. An earlier study comparing forest and grassland areas found that elements of the local environment likely drive aerobiome structure. But we wanted to know how the airborne bacterial communities varied over vertical space because, due to height differences, children may be exposed to different microbes than adults," Robinson says.

He and his team chose an urban park in Adelaide, Australia, that consists of about 700 hectares of gardens, woodlands, and playing fields. The researchers set up sampling stations in three randomly selected plots of scrub habitat, defined as parkland with semi-mature trees. At each station, they constructed a stand resembling a hat rack with petri dishes mounted at ground level and at



Vertical stratification sampling stations captured a snapshot of bacterial diversity at different heights from the ground. Image: Robinson et al. (2020); DOI: 10.1289/EHP7807.

0.5, 1, and 2 m above the ground. These levels were chosen to roughly represent the height of a jogging stroller, a 4-year-old child, and an adult, respectively.

On three different days the team left the petri dishes open for 6–8 hours to passively sample the aerobiome. They also collected soil samples. The environmental conditions during the sampling were warm and dry.

Back in the lab, the researchers extracted DNA from the samples and amplified the region coding for the 16S rRNA gene. This gene is present across all bacteria and can be used to distinguish between different organisms. ¹⁰ Software tools clustered the millions of gene sequences and taxonomically identified sequences down to the level of bacterial genus. To quantify bacterial diversity, the team used the Shannon Diversity Index, which uses both species richness (the number of species in a population) and species evenness (the abundance of each species in a population). ¹¹

The index scores declined by roughly one-third from soil level to 2 m above ground. The highest score was found in one of the soil samples. Genera that occurred in both air and soil tended to decline with height above ground. Approximately 84% of taxa in the lower air samples and 76% of upper air taxa were also found in soil. In addition, the community composition of bacterial genera varied widely at different heights. The investigators estimated that sampling height explained 22% of the variation in aerobiome makeup in the study.

Notably, this study sampled a small area for only 3 days and did not analyze bacterial DNA down to the level of species and strain. That information would be important "in order to determine whether the organisms they were looking at are the ones that might end up in our guts," Rook says. In addition, aerobiomes vary across habitat types, different weather conditions (e.g., windy vs. calm days), and concurrent activity levels (e.g., kicking a ball vs. sitting).

"The finding that microbial communities vary with height is important because we breathe hundreds of millions of microbes every day," says Emily Flies, a research fellow in health ecology at the University of Tasmania in Australia. "That exposure during early development is crucial to children's immune systems. But most aerobiome sampling is done at two meters above ground, so this study indicates that if we want to research what kids are breathing, we need to be sampling at half a meter above ground."

Flies adds that, with more research on how microbial exposure translates to health, investigators may find ways to optimize human exposure to health-promoting biodiverse microbes in greenspaces.

This study suggests that greenspaces where people can sit or lie on the ground might facilitate more biodiverse microbial exposures, she says, but further research is needed to confirm these findings and to determine whether and how those microbial exposures affect human health.

Janet L. Pelley, MS, based in Victoria, BC, Canada, also writes for Chemical & Engineering News and Frontiers in Ecology and the Environment.

References

- Flies EJ, Jones P, Buettel JC, Brook BW. 2020. Compromised ecosystem services from urban aerial microbiomes: a review of impacts on human immune function. Front Ecol Evol 8:568902, https://doi.org/10.3389/fevo.2020.568902.
- Mhuireach G, Johnson BR, Altrichter AE, Ladau J, Meadow JF, Pollard KS, et al. 2016. Urban greenness influences airborne bacterial community composition. Sci Total Environ 571:680–687, PMID: 27418518, https://doi.org/10.1016/j. scitoteny.2016.07.037.
- Robinson JM, Cando-Dumancela C, Liddicoat C, Weinstein P, Cameron R, Breed MF. 2020. Vertical stratification in urban green space aerobiomes. Environ Health Perspect 128(11):117008, PMID: 33236934, https://doi.org/10.1289/EHP7807.
- Rook GA. 2013. Regulation of the immune system by biodiversity from the natural environment: an ecosystem service essential to health. Proc Natl Acad Sci USA 110(46):18360–18367, PMID: 24154724, https://doi.org/10.1073/pnas.1313731110.
- Renz H, Skevaki C. 2020. Early life microbial exposures and allergy risks: opportunities for prevention. Nat Rev Immunol. Preprint posted online 11 September 2020, PMID: 32918062, https://doi.org/10.1038/s41577-020-00420-y.
- Kowalski K, Mulak A. 2019. Brain-gut-microbiota axis in Alzheimer's disease. J Neurogastroenterol Motil 25(1):48–60, PMID: 30646475, https://doi.org/10.5056/ inm18087.
- Aschard H, Laville V, Tchetgen Tchetgen E, Knights D, Imhann F, Seksik P, et al. 2019. Genetic effects on the commensal microbiota in inflammatory bowel disease patients. PLoS Genet 15(3):e1008018, PMID: 30849075, https://doi.org/10. 1371/journal.pgen.1008018.
- Selway CA, Mills JG, Weinstein P, Skelly C, Yadav S, Lowe A, et al. 2020. Transfer of environmental microbes to the skin and respiratory tract of humans after urban green space exposure. Environ Int 145:106084, PMID: 32977191, https://doi.org/10.1016/j.envint.2020.106084.
- Mhuireach GÁ, Betancourt-Román CM, Green JL, Johnson BR. 2019. Spatiotemporal controls on the urban aerobiome. Front Ecol Evol 7:43, https://doi.org/10.3389/fevo.2019.00043.
- Fuks G, Elgart M, Amir A, Zeisel A, Turnbaugh PJ, Soen Y, et al. 2018. Combining 16S rRNA gene variable regions enables high-resolution microbial community profiling. Microbiome 6(1):17, PMID: 29373999, https://doi.org/10. 1186/s40168-017-0396-x.
- Kim BR, Shin J, Guevarra RB, Lee JH, Kim DW, Seol KH, et al. 2017. Deciphering diversity indices for a better understanding of microbial communities. J Microbiol Biotechnol 27(12):2089–2093, PMID: 29032640, https://doi.org/10.4014/imb.1709.09027.